OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/888,182A

DATE: 05/07/2002
TIME: 14:50:40

Input Set : A:\EP.txt
Output Set: N:\CRF3\05072002\I888182A.raw

ENTERED

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       3 <110> APPLICANT: Vertex Pharmaceuticals Incorporated
       5 <120> TITLE OF INVENTION: ERK-5 DEFICIENT ANIMALS AND METHODS OF INHIBITING
ANGIOGENESIS THROUGH
       6        THE INHIBITION OF ERK-5
       8 <130> FILE REFERENCE: VPI/00-115
C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/888,182A
      11 <141> CURRENT FILING DATE: 2001-06-22
      13 <150> PRIOR APPLICATION NUMBER: US60/214,044
      14 <151> PRIOR FILING DATE: 2000-06-23
      16 <160> NUMBER OF SEQ ID NOS: 6
      18 <170> SOFTWARE: PatentIn version 3.0
      20 <210> SEQ ID NO: 1
      21 <211> LENGTH: 26
      22 <212> TYPE: DNA
      23 <213> ORGANISM: Mus musculus
      25 <400> SEQUENCE: 1
      26 cagccattcg atgtgggccc acgcta                                      26
      29 <210> SEQ ID NO: 2
      30 <211> LENGTH: 25
      31 <212> TYPE: DNA
      32 <213> ORGANISM: Mus musculus
      34 <400> SEQUENCE: 2
      35 tataacattc tcatggcgga atcgc                                       25
      38 <210> SEQ ID NO: 3
      39 <211> LENGTH: 802
      40 <212> TYPE: DNA
      41 <213> ORGANISM: Mus musculus
      43 <220> FEATURE:
      44 <221> NAME/KEY: misc_difference
      45 <222> LOCATION: (1)..(802)
      46 <223> OTHER INFORMATION: Wherein n is selected from any nucleic acid
      49 <400> SEQUENCE: 3
W--> 50 cggnacctac tgtgccctat ggaggaattc agatctgtgt aagggagtgg gccaggagga     60
      52 ggagacacag tcgggatcag cttagaagcc caggttcagt aatactgaag ttctggcagg    120
      54 gcggttgaac ccagagtgat gcgggctgtg agtccaggac attggtaggg acagttctta    180
      56 tctctcaaga gggcaagggc tggggatgtc gatcactggt aggctgatga gcatctttga    240
      58 ggttttaggt tgactctcct gtacaaaagg ggaaaagaat caagaggatt tacctcttta    300
      60 tggtcatgcc acctttggtt atatcataag ttcaaggcta gtctagaccc tgttccaaaa    360
W--> 62 gacaaaacan aaaaccnaaa cagcaatnta nganaaggga gagagggcnc agacngnccg     420
      64 ggacagatcc aaattgtaag acaacggaca caatacattg tagtgtcaca cagcagtgtc    480
      66 ctcatggcag acaactaatt attcacagaa tacctcctta aaaatagagt cttcaacata    540
      68 gcttttttcag tagctgttgg caaactgtag agtttgctct aaaattaacc atactggcca    600
W--> 70 atcttggtag atttgaatat ttctataaaa aaaattttttt ttgacagaaa ttangtccat    660
W--> 72 ggagaaagtg atttgtcaga aagcttgtaa aaaagtttgg ggctnggaaa aaacccgatt    720
```

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```
W--> 74 cggtgattaa gatcactcga tcttttaaaa gggacttggc tttaantncc ataatggnct   780
W--> 76 ttcaccgggg ggcntaaact tt                                             802
     79 <210> SEQ ID NO: 4
     80 <211> LENGTH: 794
     81 <212> TYPE: DNA
     82 <213> ORGANISM: Mus musculus
     84 <220> FEATURE:
     85 <221> NAME/KEY: misc_difference
     86 <222> LOCATION: (1)..(794)
     87 <223> OTHER INFORMATION: Wherein n is selected from any nucleic acid
     90 <400> SEQUENCE: 4
W--> 91 gattnaagat cccctcgatn tttnaaaagg acttggnttc aagggaanag ngtnttnncg    60
W--> 93 ggggnnaact tgaattggga cnccggtgtt gggatcanac tccctctttn ngcctctgta   120
W--> 95 nacccagggc acccaagtag tacacatacg ttcaggaaan catacacata cgtttaagaa   180
W--> 97 aactttataa aagttgtggc cagncggtgg tggcgcatgc ctttaatccc agcactgggg   240
     99 aggcagaggc aggcagatct cttgagtttt gggtttgagg ccaacctggt ctacaagagc   300
    101 aagcaagttc caggccagat aaggctacac agacatcttg tcttgaaaaa aagaaagaaa   360
W--> 103 gaatgaaagt tgtagaaaac ctaaaacccg gtgnnnaant ccncncttcc catgntgtta   420
    105 gtcctttggg gtttcactgt aaggccataa cctcaggaat tgggagtgcc aggggacgga   480
    107 gtgccagggg gggcttctcc ctgtgatgtg aggaggctag ctcacccgtt tcttcccatt   540
    109 ttcagctatg tggtactgga cctcatggag agcgacctac accagatcat tcactcttca   600
    111 cagccgctca ccctggaaca tgtgagatac ttcctgtacc agctgcttcg gggcctcaaa   660
    113 tacatgcact ctgctcaggt catccaccgt gatcttaaac cctctaacct tctggtcaat   720
    115 gagaactgtg agctcaagat cggtgacttt ggaatggccc gtggcctctg tacttccccc   780
    117 tgccgagcac caga                                                      794
    120 <210> SEQ ID NO: 5
    121 <211> LENGTH: 632
    122 <212> TYPE: DNA
    123 <213> ORGANISM: Mus musculus
    125 <220> FEATURE:
    126 <221> NAME/KEY: misc_difference
    127 <222> LOCATION: (1)..(632)
    128 <223> OTHER INFORMATION: Wherein n is selected from any nucleic acid
    131 <400> SEQUENCE: 5
W--> 132 caatcacggg tntnagntca ggntcaagca tggtgccaat gntgagaggc nactccatgg    60
    134 cacagttcca cttgagccca gggatggcat ctcaacatct ggacacacag gctcactagc   120
W--> 136 cacaggctgc ananaagntg gaacgnattg ttgncgaatg ccctccngtc gtgcatgaaa   180
    138 gtcttcattc tcagccacaa tggcctcctt aatgcgctcc ctggtaaggg cttcacgggt   240
    140 caaaagcaaa gtcaaaaggt ggggcgcaat caggctcatc atcagggtca tggtacttag   300
W--> 142 ccagaagggg tgcgaaggca gcagcagtna gattcgggcn ctgggttcaa ntcgcaccat   360
    144 gcgtcccagc agggagaggg cctggcggtc agcacctggg tatactgtct cccaaggcac   420
    146 aggttgcctt ggtggcaggc tctggatata ggctcgcacc ctttcagccc ccacagcctg   480
    148 aatcacagct ggtgacggag ttcccaacac catcatgatc agctgtaact ggtgcacgta   540
    150 gtttttgcct gggaagagct ggcgccgagc cagcatctca ccaaagatgc agcccacaga   600
    152 ccagaggtcg attgcctgcg gtatactcgt gc                                  632
    155 <210> SEQ ID NO: 6
    156 <211> LENGTH: 617
    157 <212> TYPE: DNA
    158 <213> ORGANISM: Mus musculus
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    160 <220> FEATURE:
    161 <221> NAME/KEY: misc_difference
    162 <222> LOCATION: (1)..(617)
    163 <223> OTHER INFORMATION: Wherein n is selected from any nucleic acid
    166 <400> SEQUENCE: 6
W--> 167 ggcaggtacc gcgttagnac cnnttatcng aacccnntgt ttntcncagn nnagcnntat    60
W--> 169 ttaaccttgn aaanagtttt tccctgaggc caagatagca natangctcn nnggagnncn   120
W--> 171 aaaaaagttt tgttctaaga ccanngaatn ggcagaatga agtggngaan gattagggag   180
W--> 173 antctggaat gacctnanta tggtgagtag gaagggaaga aggatcagtt aatncagtca   240
W--> 175 caancnnntg ctaactaacg ngcctcctnt ttatgtaagc nattagcanc ngtttcnnga   300
W--> 177 ggcagttgga aattaaaatn ttgatatatg ttacacacag ggccntgcac cacagtaggg   360
W--> 179 acttnatgnn ntntgggntc cagaagagca gtgctgaagg gacctgcagc taacttgaag   420
    181 gtactctctg gtatatgccc ttttcctgct ccccaggcca gcaggtggcc atcaagaaga   480
    183 tacctaatgc ttttgatgtg gtgaccaatg ccaaacggac cctcagggag ctgaagatcc   540
    185 tcaaacactt caaacacgac aatatcatcg ccatcaagga catcctgaag cctactgtgc   600
    187 cctatggaga attcttc                                                   617
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RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 05/07/2002
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220>
to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 4,370,377,388,391,394,409,415,417,654,705,766,768,778,794
Seq#:4; N Pos. 5,20,24,37,48,51,54,57,58,65,66,82,98,110,111,121,160,204
Seq#:4; N Pos. 394,395,396,399,403,405,415
Seq#:5; N Pos. 12,14,17,23,42,51,132,134,138,146,154,167,329,340,351
Seq#:6; N Pos. 18,22,23,29,36,37,43,46,50,51,52,56,57,70,74,101,105,110,111
Seq#:6; N Pos. 112,117,118,120,144,145,150,166,170,182,196,198,234,244,246
Seq#:6; N Pos. 247,248,261,269,281,289,291,297,298,320,345,365,369,370,371
Seq#:6; N Pos. 373,378